Coronary Artery Risk Development in Young Adults (CARDIA) dbGaP Acknowledgment Statements Revised on 12-01-2012

The acknowledgment of the CARDIA dataset(s) should always include acknowledgment of the CARDIA contract (component I below) **AND** the genotypic/genomic dataset(s) (component II below). In addition, if the CARDIA phenotypic dataset(s) used in your analysis were derived from non-contract supported studies, these dataset(s) should also be acknowledged (component III below).

Components of the CARDIA dbGaP Acknowledgment Statements:

I. Coronary Artery Risk Development in Young Adults (CARDIA) Contract Acknowledgment (Use the following statement for ALL analyses resulting from CARDIA dbGaP data).

"The Coronary Artery Risk Development in Young Adults Study (CARDIA) is conducted and supported by the National Heart, Lung, and Blood Institute (NHLBI) in collaboration with the University of Alabama at Birmingham (N01-HC95095 & N01-HC48047), University of Minnesota (N01-HC48048), Northwestern University (N01-HC48049), and Kaiser Foundation Research Institute (N01-HC48050). This manuscript was not approved by CARDIA. The opinions and conclusions contained in this publication are solely those of the authors, and are not endorsed by CARDIA or the NHLBI and should not be assumed to reflect the opinions or conclusions of either."

- **II. Genotypic/Genomic Dataset Acknowledgment** (Use one or more of the following statements relevant to the specific datasets used in your analysis)
 - **A. Candidate gene Association Resource (CARe):** "Funding for CARe genotyping was provided by NHLBI Contract N01-HC-65226."
 - **B.** The Genes, Environment Association Studies consortium (GENEVA): "Genotyping for the CARDIA GENEVA cohort was supported by grant U01 HG004729 from the National Human Genome Research Institute."
 - C. NHLBI Grand Opportunity Exome Sequencing Project (GO-ESP): "This study is part of the NHLBI Grand Opportunity Exome Sequencing Project (GO-ESP). Funding for GO-ESP was provided by NHLBI grants RC2 HL103010 (HeartGO), RC2 HL102923 (LungGO) and RC2 HL102924 (WHISP). The exome sequencing was performed through NHLBI grants RC2 HL102925 (BroadGO) and RC2 HL102926 (SeattleGO).

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(FHS), the Jackson Heart Study (JHS) and the Multi-Ethnic Study of Atherosclerosis (MESA)."